

Replace the paragraph beginning at page 8, line 1, with the following rewritten paragraph:

--For example, human TRANCE antisense oligonucleotides that consist of 24 to 50 nucleotides encompassing the translation initiation codon can be used to inhibit TRANCE activity. In Homo sapiens tumor necrosis factor (ligand) superfamily, member 11 (TNFSF11), mRNA splice variant 1 (GenBank Accession # NM_003701.2), this would include the region from nucleotides 141 to 190 (SEQ ID NO:17), including the start codon at nucleotide number 157 (see FIG. 1). In mRNA splice variant 2 (GenBank Accession # NM_033012.1), this would include nucleotides 81 to 140 (SEQ ID NO:18), including the start codon at nucleotide number 95 (see FIG. 3).--

In the drawings:

Please substitute new Figure 2 for the original Figure 2.